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# RNA *in situ* hybridization of dissected gonads\*

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## 1. Introduction

Determination of the temporal and spatial pattern of gene expression is an important step in understanding gene function during *C. elegans* development. The expression pattern of a gene can be examined in several ways including transgenic reporter fusions (e.g., gfp,  $\beta$ -galactosidase), as well as immunohistochemistry and RNA *in situ* hybridization for analysis of the endogenous protein and mRNA, respectively. While reporter fusions and immunohistochemistry require time and effort to generate reagents (reporter construction and generation of transgenic lines or antibody production), RNA *in situ* hybridization only requires the transcribed region of a gene of interest. In addition, as a complement to protein expression patterns of a gene, RNA *in situ* allows one to determine whether the mRNA is subjected to post-transcriptional controls such as differential RNA localization, stability, and/or translational regulation (Evans et al., 1994; Seydoux and Fire 1994; Jones et al., 1996; Lee and Schedl 2001, 2004).

Several *in situ* hybridization protocols have been developed to detect RNA in *C. elegans*. RNA *in situ* in embryos is well described previously (Seydoux and Fire, 1995). Protocols for whole-mount *in situ* of intact animals and embryos can be also found at Yuji Kohara's website. Fluorescence RNA *in situ* hybridizations have been also described (Albertson et al., 1995; Graves, et al., 1999; Pitt et al., 2000). In this chapter, a protocol for detection of mRNA by *in situ* hybridization to dissected gonads is described. This protocol is primarily based on the method of Seydoux and Fire (1995) as adapted for dissected gonads.

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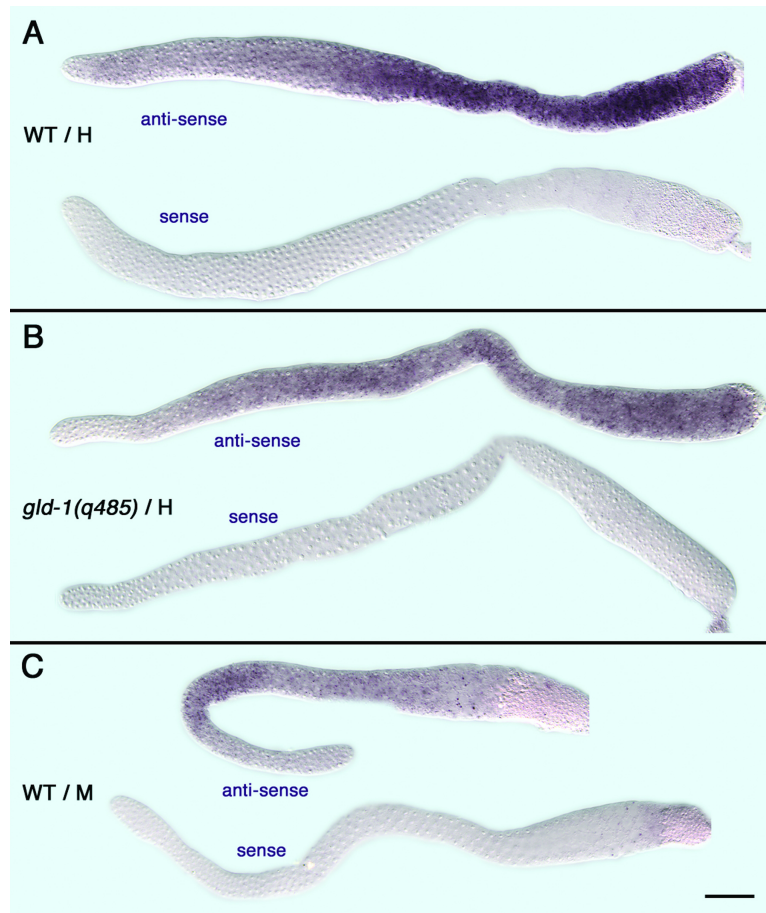
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In contrast to an earlier method described for dissected gonads wherein the gonads were affixed to slides (Crittenden et al., 1994), the various manipulations described below (gonad dissections, hybridizations, and washes) are all performed in solution, either in glass dishes (dissection) or in small glass tubes (hybridization). Washes are done by low-speed spins and all solutions (except fixatives) contain 0.1% Tween 20 to prevent dissected gonads from sticking to glass. A key to make this protocol work consistently is to fix dissected gonads in a combination of paraformaldehyde (3%) and glutaraldehyde (0.25%). The addition of glutaraldehyde results in a lower background, as compared to fixation with paraformaldehyde alone and makes gonads both less sticky and less susceptible to breakage.

It is important to perform RNA *in situ* hybridization using both sense and anti-sense probes for a gene of interest, at least for initial experiments. RNA *in situ* hybridization with a sense probe should give little or no signal compared to that with an anti-sense probe (Figure 1). The sense probe background control is particularly important for detecting transcripts that are expressed at very low levels, which requires an extended developing time with the alkaline phosphatase-mediated detection system. In general, we can easily detect most mRNAs that have five or more EST clones reported in WormBase. For mRNAs that have less than two EST clones (therefore likely expressed at very low levels), it is helpful to concentrate the probe either by using more template in the asymmetric PCR (probe synthesis step) or by dissolving the digoxigenin (DIG) labeled probe in a smaller volume. RNA *in situ* hybridization with a concentrated anti-sense probe usually results in increased signal with decreased developing time of the alkaline phosphatase-mediated detection system. In addition, concentration of the sense probe does not increase the background signal significantly.



**Figure 1. Gemline RNA *in situ* hybridization with an anti-sense and a sense probes.** Representative photomicrographs of RNA *in situ* hybridizations showing *cep-1/p53* mRNA accumulation in wild-type (A) and *gld-1*(null) (B) hermaphrodites, and wild-type male (C). Each panel has images from gonads hybridized with an anti-sense probe and a sense probe otherwise treated equally. Each image shows a surface view of the gonad where the distal region is on the left and the proximal region is on the right. Sense probe gave no or very little signals (the proximal signal is likely due to endogenous phosphatases). In wild-type hermaphrodites, *cep-1* mRNA accumulates throughout the germ line and its levels increase toward the proximal region of the germ line. In *gld-1*(null), *cep-1* mRNA levels are slightly lower as compared with wild type. In wild-type males, *cep-1* mRNA accumulates at higher levels only in the distal meiotic germ cells, showing sexual dimorphism.

## 2. General expression pattern

The levels of most maternal mRNAs become higher in the middle of the pachytene region and remain high in the developing oocytes in the proximal region because the bulk of RNA synthesis for oogenesis occurs in pachytene stage germ cells, with progressively less RNA synthesis as germ cells proceed through the diplotene stage. As chromosomes become highly condensed in late stage oocytes (diakinesis), almost no *de novo* RNA synthesis occurs (Gibert et al., 1984; Schisa et al., 2001) and RNA polymerase II becomes inactive (Kelly et al., 2002). Therefore, most maternal mRNAs are transcribed primarily in pachytene stage germ cells in the distal region and transported into the common cytoplasmic core. In the proximal region, maternal mRNAs are packaged into growing oocytes. These processes result in extensive RNA accumulation throughout pachytene stage germ cells and developing oocytes (Figure 1).

Some mRNAs are expressed throughout the germline including the distal mitotic region (e.g., *glp-1*) while others are only expressed in meiotic cells (e.g., *rme-2*). For a number of transcripts expressed throughout the germline, the relative levels observed in the distal mitotic region are lower, compared to the levels in the meiotic pachytene cells and proximal oocytes (Figure 1). The difference is likely, at least in part, a reflection of a lower level of the mRNA/protein being required for function in the distal mitotic region, while a higher level is required for late oogenesis and/or maternally for early embryogenesis.

## 3. Materials

Solutions do not need to be treated with diethyl pyrocarbonate.

PBS: Make a 10x stock solution according to Sambrook et al. (1989).

PBTw: 1x PBS containing 0.1% Tween 20.

Paraformaldehyde / glutaraldehyde fix: Paraformaldehyde/Glutaraldehyde fixation works well for germline RNA *in situ*: gonads remain intact through an extensive protease treatment (50 µg/ml Proteinase K for 30 min at room temperature (RT) that would destroy gonads fixed in paraformaldehyde alone. This fixation can also be used with good results for histological staining with some antibodies.

To make 3% paraformaldehyde / 0.25% glutaraldehyde / 0.1M K<sub>2</sub>HPO<sub>4</sub> (pH7.2), mix

1. 10 ml 16% formaldehyde (sealed ampoules from EM Sciences #15710)
2. 0.53 ml 25% glutaraldehyde (sealed ampoules from EM Sciences #16220)
3. 25 ml 0.2 M K<sub>2</sub>HPO<sub>4</sub> (pH7.2)

and dispense in 10 ml aliquots and store at -20°C.

Proteinase K: 20 mg/ml. Dispense in 10 µl aliquots and store at -80°C. Do not refreeze, discard after use.

Hybridization buffer: 5X SSC, 50 % deionized formamide, 100 µg/ml autoclaved Herring sperm DNA, 50 µg/ml Heparin and 0.1 % Tween-20. We make 50 ml, dispense in 10 ml aliquots and store at -20°C.

Alkaline-phosphatase-conjugated anti-DIG (Fab2 fragment) from Roche (#1 093 274): An optimal dilution of this antibody needs to be determined empirically as lot-to-lot variations may exist. We have been using dilutions from 1:1000 to 1:2500.

Staining Solution: 100 mM NaCl; 5 mM MgCl<sub>2</sub>; 100 mM Tris (pH 9.5); 0.1% Tween 20; 1mM Levamisole. Levamisole is a potent inhibitor of endogenous phosphatases.

Sigma Fast BCIP/NBT tablet (#B5655): Dissolve one tablet in 10 ml staining solution. If non-specific background signal is high, you may want to use more staining solution (up to 20 ml) to dissolve the tablet.

Anti-Fade solution: Make 0.2M Dabco (Sigma #D2522) in 20 mM Tris (pH 8.0), which can be stored at -20°C. Then dilute this solution (10%) in glycerol (90%) and store -20°C.

Culture tubes: Borosilicate glass tubes with plain end. These tubes can be baked to decontaminate RNase. Available from Fisher as 6 mm (O.D.) x 50 mm (length) in either Pyrex (Fisher #14-957AA) or Kimax (Fisher #14-925B) brands.

Dissection dishes: We use Square Watch Glass from Carolina Biological Supply (#ER-74-2300).

QIAquick PCR Purification kit from Qiagen (#28104): Works well to remove unincorporated dNTP and PCR primers.

DIG DNA Labeling Mix, 10 x conc.: Roche (#1 277 065).

## 4. Procedure

Single-stranded probes from cloned cDNA can be prepared anytime. Following dissections and fixation, fixed gonads can be stored in 100% MeOH at  $-20^{\circ}\text{C}$  for at least one week. Once the gonads are subjected to permeabilization, the remaining procedure has to be completed.

Procedure 1. Preparation of single-stranded probes from cloned cDNA

Based on Patel and Goodman (1992), with slight modifications.

### 1. PCR amplification of a cDNA of interest with any set of primers.

A cloned cDNA template is optimal because amplified products from genomic DNA usually generate high background.

Use less than 1  $\mu\text{l}$  (10 pmole/ $\mu\text{l}$ ) of primers because QIAquick PCR Purification kit will not effectively remove primers if the concentration is higher.

We usually set up two 50  $\mu\text{l}$  PCR reaction with 35 cycles ( $94^{\circ}\text{C}$  30 seconds,  $55^{\circ}\text{C}$  30 seconds and  $72^{\circ}\text{C}$ , 1 minute/1 kb).

10x Taq Buffer	5.0 $\mu\text{l}$
dNTP mix (2.5 mM each)	2.0 $\mu\text{l}$
5'Primer (10 pmole/ $\mu\text{l}$ )	1.0 $\mu\text{l}$
3'Primer (10 pmole/ $\mu\text{l}$ )	1.0 $\mu\text{l}$
a cloned cDNA (1:50 dilution of miniprep DNA) or 5–10 ng of purified plasmid	2.0 $\mu\text{l}$
Taq Polymerase	2 units
ddH <sub>2</sub> O	to 50 $\mu\text{l}$

Run 3  $\mu\text{l}$  each in an agarose gel; the PCR product must be an intense single band.

### 2. Purification of PCR product

Use the QIAquick PCR Purification kit to purify the PCR product away from unincorporated dNTP and PCR primers. Elute with 25  $\mu\text{l}$  Elution buffer. The concentration should be above 200 ng/ $\mu\text{l}$ .

### 3. Asymmetric (one-way) PCR

To generate DIG labeled probe (35 cycles,  $94^{\circ}\text{C}$  30 seconds,  $55^{\circ}\text{C}$  30 seconds and  $72^{\circ}\text{C}$ , 1 minute/1 kb):

10x Taq Buffer	2.5 $\mu\text{l}$
DIG DNA Labeling Mix, 10 x conc	2.5 $\mu\text{l}$
Primer <sup>‡</sup> (10 pmole/ $\mu\text{l}$ )	1.0 $\mu\text{l}$
Purified PCR product (at least 200 ng/ $\mu\text{l}$ )	4.0 $\mu\text{l}$

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<sup>‡</sup>Use only one primer. We set up two reactions, one to generate a sense probe with 5'primer and the other to generate an anti-sense probe with 3'primer. RNA *in situ* hybridization with the sense probe will be a background control.

Taq Polymerase	2 units
ddH <sub>2</sub> O	to 25 µl

4. 75 µl of H<sub>2</sub>O is added to the reaction and 90–95 µl of the diluted reaction is transferred to a new tube.
5. 10 µl of 1M NaCl and 3 vols of 100% EtOH are added to the diluted reaction. After 30 min at –70°C or overnight at –20°C, the reaction is centrifuged at 14,000 rpm for 10 min. The pellet is washed in 70% ethanol, dried, and resuspended in 400 µl of hybridization buffer.
6. The probe is boiled for 1 hr. This step reduces the length of the probe for efficient penetration.
7. Probes can be stored at –20°C for a long period. We have used two year-old probe without any obvious reduction of signal.

#### Procedure 2. Dissections and fixation

1. Wear latex gloves (optional), prepare dissected gonads (see [http://www.genetics.wustl.edu/tslab/gonad\\_dissections.html](http://www.genetics.wustl.edu/tslab/gonad_dissections.html) for dissection) and fix in 3% paraformaldehyde / 0.25% glutaraldehyde / 0.1 M K<sub>2</sub>HPO<sub>4</sub> (pH7.2) for 1.5 to 2 hr at RT. We routinely dissect about 150 animals per each hybridization to obtain 40 or more gonads that are completely extruded, well stained, and unbroken at the end of the entire procedure.
2. Transfer entire contents to a 5 ml glass conical tube and spin in clinical table top centrifuge for 1 min at setting 3 (All subsequent spins are also for 1 min at setting 3).
3. Discard the fixative in the appropriate chemical waste container and resuspend in 2–4 ml PBTw.
4. Spin and discard PBTw as much as possible.
5. Add about 2 ml of 100% cold (–20°C) Methanol (MeOH). At this point, dissected gonads can be stored in 100% MeOH at –20°C for at least one week with no obvious problems.

#### Procedure 3. Permeabilization: Proteinase K digestion

1. Add 2–3 ml of PBTw to the MeOH solution, and spin. Wash twice in PBTw. Now split the sample into 2 or more tubes (either into 5 ml conical tubes or into 6 mm x 50 mm culture tubes).
2. Using a 20 mg/ml stock of Proteinase K (PrK), make solutions of 50 µg/ml PrK in PBTw. Add 2ml PrK / PBTw and digest for 30 min at RT.
3. Wash three times in PBTw.
4. Fix 15 min in 3% paraformaldehyde / 0.25% glutaraldehyde / 0.1M K<sub>2</sub>HPO<sub>4</sub> (pH7.2)
5. Incubate 15 min in PBTw containing 2 mg/ml glycine.
6. Wash three times in PBTw.
7. Divide gonads among several 6 x 50 mm culture tubes, one tube per hybridization.

#### Procedure 4. Hybridization and wash

1. Place gonads in 400 µl of 50% PBTw / 50% hybridization buffer (HB) and incubate 5 min at 48°C (in water bath).
2. Pre-hybridize in 400 µl HB for 1 hr at 48°C.
3. Dilute boiled probe in HB to give 100 µl of solution. Try a 1:2 dilution for probes to messages of unknown abundance. Experience with *gld-1* probes suggests that probe dilution is not critical – a strong, clean signal was

obtained at dilutions between 1:2 and 1:5, but it began to fall off at 1:10. Probe to major sperm protein RNA, which is very abundant during spermatogenesis, could be diluted 1:20 without any fall off in signal.

4. Boil diluted probe 5 min, cool to 48°, and add to gonads.
5. Hybridize for 12 to 36 hr at 48°C (longer hybridization times tend to give increased signal and lower background). We routinely hybridize for 36 hrs.
6. Wash three times in PBTw.
7. Divide gonads among several 6 x 50 mm culture tubes, one tube per hybridization.
8. Add 400 µl of HB that is pre-warmed to 48°, spin down, and remove supernatant.
9. Divide gonads among several 6 x 50 mm culture tubes, one tube per hybridization.
10. Wash 4 x 15 min in HB at 48°C.
11. Then perform the following washes:
  - a. 1 × 10 min in 50% HB / 50% PBTw at 48°
  - b. 2 × 10 min in PBTw at 48°
  - c. 1 × 5 min in PBTw at RT

#### Procedure 5. Probe detection

1. Incubate 15 min in PBTw /0.5 mg/ml BSA (either fraction V or restriction enzyme grade).
2. Dilute alkaline-phosphatase-conjugated anti-DIG (Fab2 fragment) to optimal conc. in PBTw /BSA. Add 500 µl to each tube and incubate overnight at 4°C or at RT for 2 hr. For gonads, overnight at 4°C is definitely better than 2 hr at RT.
3. Wash at RT as follows:
  - a. 2 x 2 min PBTw
  - b. 3 x 20 min PBTw / BSA
  - c. 2 x 5 min in Staining solution.
4. Transfer gonads to staining solution with BCIP/NDT and 100 ng/ml DAPI and cover to protect from light. Signal will come up anywhere between 5 min and 6 hr. For convenience, the reaction can be monitored by transferring tube contents to a glass dish and inspecting periodically under the dissecting scope. However, stain will appear darker in the dissecting scope than under Nomarski optics with a 40X or higher lens.
5. Stop the reaction by washing three times in PBTw. Finally place gonads in PBS containing 100 ng/ml DAPI.
6. After removing DAPI solution as much as possible, add 35 µl anti-fade solution.

#### Procedure 6. Mounting gonads for viewing

- Using a drawn capillary pipette, transfer settled worms onto a large 2% agar pad that covers most of a slide. After drawing off excess liquid with a capillary, an eyelash hair can be used to push gonads and intestines away from one another. Cover with a large (24 x 50 mm) coverslip, taking care not to move the coverslip once in place. Also do not seal the coverslip immediately – image may improve as liquid evaporates and gonads become somewhat flattened. We usually take images after storing the slides at 4°C overnight without sealing.



Bright field (Nomarski) images must be taken before DAPI images because the strong UV light focused by the microscope literally burns gonads brown. Slides can be stored at 4°C for a week or more, particularly if sealed with nail polish around the periphery of the coverslip. Because the stain is purplish, images are better in color than in black and white.

## 5. Acknowledgements

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## 6. References

- Albertson, D.G., Fishpool, R.M., and Birchall, P.S. (1995). Fluorescence *in situ* hybridization for the detection of DNA and RNA. *Methods Cell Biol.* 48, 339–364. [Abstract](#)
- Crittenden, S.L., Troemel, E.R., Evans, T.C., and Kimble, J. (1994). **GLP-1** is localized to the mitotic region of the *C. elegans* germ line. *Development* 120, 2901–2911. [Abstract](#)
- Evans, T.C., Crittenden, S.L., Kodoyianni, V., and Kimble, J. (1994). Translational control of maternal *glp-1* mRNA establishes an asymmetry in the *C. elegans* embryo. *Cell* 77, 183–194. [Abstract Article](#)
- Gibert, M.A., Starck, J., and Beguet, B. (1984). Role of the gonad cytoplasmic core during oogenesis of the nematode *Caenorhabditis elegans*. *Biol. Cell* 50, 77–85. [Abstract](#)
- Graves, L.E., Segal, S., and Goodwin, E.B. (1999). **TRA-1** regulates the cellular distribution of the *tra-2* mRNA in *C. elegans*. *Nature* 399, 802–805. [Abstract Article](#)
- Jones, A.R., Francis, R., and Schedl, T. (1996). **GLD-1**, a cytoplasmic protein essential for oocyte differentiation, shows stage- and sex-specific expression during *Caenorhabditis elegans* germline development. *Dev. Biol.* 180, 165–183. [Abstract Article](#)
- Kelly, W.G., Schaner, C.E., Dernburg, A.F., Lee, M.H., Kim, S.K., Villeneuve, A.M., and Reinke, V. (2002). X-chromosome silencing in the germline of *C. elegans*. *Development* 129, 479–492. [Abstract](#)
- Lee, M.H., and Schedl, T. (2001). Identification of *in vivo* mRNA targets of **GLD-1**, a maxi-KH motif containing protein required for *C. elegans* germ cell development. *Genes Dev.* 15, 2408–2420. [Abstract Article](#)
- Lee, M.H., and Schedl, T. (2004). Translation repression by **GLD-1** protects its mRNA targets from nonsense-mediated mRNA decay in *C. elegans*. *Genes Dev.* 18, 1047–1059. [Abstract Article](#)
- Patel, N.H., and Goodman, C.S. (1992). Preparation of digoxigenin-labeled single-stranded DNA probes. In: *Non-Radioactive Labeling and Detection of Biomolecules*, C. Kessler, ed. (Berlin: Springer-Verlag).
- Pitt, J.N., Schisa, J.A., and Priess, J.R. (2000). P granules in the germ cells of *Caenorhabditis elegans* adults are associated with clusters of nuclear pores and contain RNA. *Dev. Biol.* 219, 315–333. [Abstract Article](#)
- Sambrook, J., Fritsch, T., and Maniatis, T. (1989). *Molecular cloning: a laboratory manual*. 2nd edition. (New York: Cold Spring Harbor Laboratory Press).
- Schisa, J.A., Pitt, J.N., and Priess, J.R. (2001). Analysis of RNA associated with P granules in germ cells of *C. elegans* adults. *Development* 128, 1287–1298. [Abstract](#)
- Seydoux, G., and Fire, A. (1994). Soma-germline asymmetry in the distributions of embryonic RNAs in *Caenorhabditis elegans*. *Development* 120, 2823–2834. [Abstract](#)
- Seydoux, G., and Fire, A. (1995). Whole-mount *in situ* hybridization for the detection of RNA in *Caenorhabditis elegans* embryos. *Methods Cell Biol.* 48, 323–337. [Abstract](#)



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